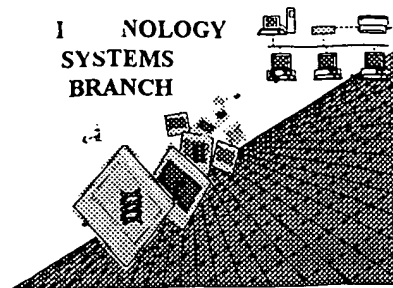


5000

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



## **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/857,372A  
Source: Pt/09  
Date Processed by STIC: 10/29/2001

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**

**<http://www.uspto.gov/web/offices/pac/checker>**

PCT09

## RAW SEQUENCE LISTING

DATE: 10/29/2001

PATENT APPLICATION: US/09/857,372A

TIME: 15:02:39

Input Set : A:\JAB-1430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

pg 3-5

**Does Not Comply  
Corrected Diskette Needed**

OK  
C-→

```

3 <110> APPLICANT: Contreras, Roland
4   Nelissen, Bart
5   DeBacker, Marianne
6   Luyten, Walter
7   Viaene, Jasmine
8   Logghe, Marc
10 <120> TITLE OF INVENTION: Drug Targets in Candida Albicans
12 <130> FILE REFERENCE: 53731/000
14 <140> CURRENT APPLICATION NUMBER: US/09/857,372A
15 <141> CURRENT FILING DATE: 2001-09-24
17 <150> PRIOR APPLICATION NUMBER: 982204122.0
18 <151> PRIOR FILING DATE: 1998-12-04
20 <160> NUMBER OF SEQ ID NOS: 22
22 <170> SOFTWARE: PatentIn Ver. 2.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 438
26 <212> TYPE: DNA
27 <213> ORGANISM: Candida albicans
29 <400> SEQUENCE: 1
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31 taactgatga acaattggaat accattgcat tgacatttgg ttttgcttca ataataattaa 120
32 tcataatata tcatgccata tctactaatg tacataaatt agaagatgaa accccatcat 180
33 cttcatttac cagaacaaat actactgaaa ctactgttgc aagtaagaaa aagaagtaat 240
34 aactgatgga tttttcttcc taccaccaat tgaataatgc tagacttggt ggtgtgctac 300
35 aaatatattca aaagaaaata cgaatacttt ataaaatggt aagaacggaa gatggtttct 360
36 catttatata ctaaatacaa atcacatata catacacaaa cacaataata tacatacacc 420
37 tatatccctt tatttgat                                     438
39 <210> SEQ ID NO: 2
40 <211> LENGTH: 1380
41 <212> TYPE: DNA
42 <213> ORGANISM: Candida albicans
44 <400> SEQUENCE: 2
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46 ccggccttga tcagaacctc cgctgccttc gaccgttcta aacctcatgt caacattggt 120
47 actattgggtc atgttgatca tggtaaaact acattgactg ctgctatcac caaagtttta 180
48 gccgaacaag gtggtgccaa cttcttggat tatggttcta ttgatagagc tccagaagaa 240
49 agagctagag gtatcactat ttccactgcc cacgttgaat acgaaaccaa gaacagacac 300
50 tatgcccacg ttgattgtcc aggacacgct gattatatca aaaatatgat tactgggtgc 360
51 gctcaaatgg atggtgctat cattgttgtt gctgccactg atgggtcaaat gcctcaaacc 420
52 agagaacatt tgttattggc cagacaagtt ggtgttcaag acttggttgt gtttgtcaac 480
53 aaagtcgata ctattgatga cctgaaatg ttggaattag tcgaaatgga aatgagagaa 540
54 ttgttatcca cctacggttt tgatggtgac aacactccag ttattatggg atctgcttta 600
55 atggcctttg aagacaagaa accagaaatt ggtaagggaag ctatcttgaa attgttagat 660
56 gctgtcgatg aacacattcc aactccatca agagacttgg aacaaccatt tttgttacca 720
57 gttgaagacg tgttctccat ctccggtaga ggaactgttg tcaactgtag agttgaaaga 780
58 ggtgttttga agaagggtga agaaatcgaa attgttggtg gttttgacaa accttacaag 840
59 actactgtta ccggtattga aatgttcaaa aaagaattag actctgctat ggctggtgac 900

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## RAW SEQUENCE LISTING

DATE: 10/29/2001

PATENT APPLICATION: US/09/857,372A

TIME: 15:02:39

Input Set : A:\JAB-1430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

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60 aactgtggtg ttttgttaag aggtgttaaa agagatgaaa tcaagagagg tatggttttg 960
61 gccaaaccag gtactgctac ttctcacaag aagttcttgg ctctcttgta tattttgact 1020
62 tccgaagaag gtggtcggtc cactccattt ggtgaagggtt acaagcctca atgcttcttc 1080
63 agaactaacg atgtcactac cacattttca ttcccagaag gagaagggtg tgatcattct 1140
64 caaatgatca tgccagggtg caacattgaa atggttggtg aattgatcaa atcttggtcca 1200
65 ttagaagtca accaacgttt caacttgaga gaagggtgta aaactgttgg tactggtttg 1260
66 attaccagaa tcatcgaaat aacagaatgt gcactgtgaa taataaaaaa aaaaagggtg 1320
67 tatatagggtg actttgtatt ttgtattgaa caataaaaatt ctgtaaatag taagggcctc 1380
69 <210> SEQ ID NO: 3
70 <211> LENGTH: 2283
71 <212> TYPE: DNA
72 <213> ORGANISM: Candida albicans
74 <400> SEQUENCE: 3
75 gaattcgccc ttaagcactc gtttcaacta tacattcagt aacaacaccc ttaatttacc 60
76 aaactacatt aatggaagta acacaacgga cgcagagtca gacacaacca acacaacagt 120
77 caccgacaac tcagacgcaa acccaaagca aagaggacca gaataggatt tgtcaattga 180
78 tttgctccac gggtcagttt ggcaattatg atttgaatat caacgataaa actatcgtac 240
79 aaggtaaaaat gacgtggtat ttggaagag accccaactc agatttgcaa gtggcgctcg 300
80 cgtcgagaat ttcaaacaaag cattttcaaa tctggctcaa cttcaatgat aaatcactat 360
81 ggataaagga cacttcaact aacgggacac accttaacaa cagtcgattg gtgaaaggat 420
82 caaactacct tcttaatcag ggtgatgaaa tagcagtagg ggttggtaga gacgaggacg 480
83 ttgtgagggtt tgtcgttgtc tttggtgaca aatacaaccc ggcaaagcta cctgattcga 540
84 ccaacacaat taaagatgaa ggaatataca aagactttat tgtgaaaaat gaaacgatag 600
85 gccaaaggagc atttgccact gtgaaaaaagg cgattgaacg atctacgggc gagtctgacg 660
86 cggatgaagat tataaatcga agaaaagcat taaataccgg tggatggaagt gccatggcag 720
87 gagtggaccg tgaattgtcc atattagagc ggctcaacca cccaaatata gttgctctaa 780
88 aagcttttta tgaagatatg gacaattact atattgtgat ggaattgggt ccgggcgggtg 840
89 atttgatgga ctttgtggct gcaaacggtg caataggaga agacgcaaca caagtgatca 900
90 cgaaacagat tctagaagga attgcctatg ttcataattt aggaatctcc catcgtgatt 960
91 tgaagccaga taatattttg attatgcaag atgacccaat acttgtaaaa atcacccgact 1020
92 ttggattggc aaaattcagt gacaatctga cgtttatgaa aactttttgt ggtacattgg 1080
93 cgtatgttgc tcccgaagtt atcacccgta agtatggatc atcgagatg gaactgcaac 1140
94 aaaaggacaa ctactcttcc ttggttgaca tttggtcttt gggatgtttg gtttatgtac 1200
95 ttttaacttc tcattttacca ttcaacggga aaaaccagca acaaatgttt gccaaagatca 1260
96 aaaggggcca atttcatgag gctccattaa attcatacga catttctgaa gacggaagag 1320
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98 tgaaacataa atggttgcaa gacttgtatg aagaggattc tgtcaaatca ttgagtttat 1440
99 cgcaatcaca gtcgcaacaa tctcgaaaga tagataatgg tatccatata gaatcattga 1500
100 gcaaaattga tgaagacgtt atgcttcgtc cattggatag cgaaagaaat aggaaatcaa 1560
101 gtaaacagca agattttcaag gtacccaagc gtgtgattcc gttatctcaa catcctgcaa 1620
102 caccgttacc aatgtcacaac ccgaaaaaga ggccgtatca aatagacctt agaacaaaca 1680
103 aaaaagtcga tttggaagaa cctctgacaa gcaagaaagt caagctaagt gattccgttg 1740
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105 atatttcaaa gtccccgttt tctttcgga gaaatgacac ttgtgattgc gagatagacg 1860
106 acgacagact atccaaactt cattgtgtca ttaccaaaga aaacgactct atatggttat 1920
107 tggataagag tactaactcg tgcttggtca acaatactag tgttggaaaa ggcaacaaaag 1980
108 ttttgcttag aggaggggag atattacatc tcttctttga cccattgtca ctgcaacata 2040
109 taggtttcaa agtagtcctt gttgatcaac tgtctggtga acataagagt caagtggagg 2100
110 ttttgaaca aacctcagaa gaaatgaata ttattccact tatttctggt ttaagttagta 2160

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## RAW SEQUENCE LISTING

DATE: 10/29/2001

PATENT APPLICATION: US/09/857,372A

TIME: 15:02:39

Input Set : A:\JAB-1430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

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111 taagttcata gatttagcat atatacaagc atttcctata gaaacaaagg ttcattaatt 2220
112 tagttattta cctccatgca attacattta cttcttcttc caagggcgaa ttctgcagat 2280
113 atc 2283

```

115 &lt;210&gt; SEQ ID NO: 4

116 &lt;211&gt; LENGTH: 826

117 &lt;212&gt; TYPE: DNA

118 &lt;213&gt; ORGANISM: Candida albicans

120 &lt;400&gt; SEQUENCE: 4

```

121 atgggtagta tgtgaagata caatattgaa agtgtttact agaatatcta agatgtttga 60
122 gcccatggac atttttggat ttgataatta aaaaaagtag caatagatta ttgcgttga 120
123 gaaagaatca ccatagtgc aagatttgat agatgtttaa atgttcacgc aggcgaaaga 180
124 tgtaacatct cttaaagtaa gaagaatatg gacatgaata aaaatagata gcaactatttt 240
125 ggaacttggt gaagatatta aaatagaatg ggatttcaac atagatattc aaagtaacga 300
126 aacctcacia tcaataaaaa acaacagtaa tactaacaat tcaattttta tttttataga 360
127 gggtagtcca tctttaggta aacgtcacaa caaatctcac accttatgta acagatgtgg 420
128 ccgtcgttca ttccacgtcc aaaagaagac ctgttcttct tgtggttacc cagctgctaa 480
129 aatgagatct cacaactggg ccttaaaaagc caaaagaaga agaactactg gtaccggtag 540
130 aatggccttac ttgaaacacg ttaccagaag attcaagaac ggtttccaaa ctggtgttgc 600
131 taaagctcaa accccttccg cttaaaactaa ttactgaagt tattggtcat gcattagtca 660
132 ttattcatta aagtcattgt aagcatagca aaggaagaat tggtagatt ctgtttaaa 720
133 atgtaatgac tatttaatat ctgtttaaat aagaggttta gtctttattt ttttacgtat 780
134 acaccaaaaa aaaaagaaac aaataaaatc tgtatattaa tgttgg 826

```

136 &lt;210&gt; SEQ ID NO: 5

137 &lt;211&gt; LENGTH: 978

138 &lt;212&gt; TYPE: DNA

139 &lt;213&gt; ORGANISM: Candida albicans

141 &lt;400&gt; SEQUENCE: 5

```

142 atgggtacta gtacaagtga agcattgaag aacatcaaaa acaaacagcg aagacagaaa 60
143 gtttttgcag aaataaaaaca tgaaaagaat aaacaacgctc ataagcaaag agccgaaaga 120
144 gctaaggagag aaagagaaaaa ccccgaaatta agagaggaaa gaatagcagc taatatcca 180
145 gatactatag atagcaaacg tattttatgat gagactatag ctgctgaagt tgaaggagat 240
146 gacgagtttc agtcataattt cactaaacttg ttggaagaac caaagatttt gttgacaaca 300
147 agtgccaatg ctaaaaaaacc ggcctatgaa tttgcagaca tgatcatgga ctttttaccg 360
148 aatgtgacat ttatcaaaaag gaagaaggaa tatacaatgc aagatatggc caaatattgc 420
149 tcgaatagag acttcactgc attgcttgtc atcaacgaag acaagaagaa ggtcaatgg 480
150 ataacgctca tcaattttacc tgaagggcc aacattttatt tttcgattac atcaatagtt 540
151 gatgggaaaa gaattaaggg acacgggaaa gctggtgatt atttacctga gattgtattg 600
152 aataatttca attcaagatt gggtaaaact gtgggaagac tatttcaaag tattttccct 660
153 cataaacctg aacttcaagg aagacaagtg attactttgc acaatcaacy tgattatatt 720
154 tttttcagaa gacatagata tattttcaga aatgaggaaa aggttggatt gcaggaaattg 780
155 ggtccgcagt ttacattaaa gctaagaaga atgcaaaagg gagtacgtgg tgatgttgtt 840
156 tgggaacaca gaccagatat ggaaagagat aagaagaagt tttatttata agcgggtgta 900
157 taaaggtagt agtagtgcgt ttataagtat gtgtgtgtgt ttatgcatag atgtgtaaa 960
158 agtaatacag ctaattcg 978

```

160 &lt;210&gt; SEQ ID NO: 6

161 &lt;211&gt; LENGTH: 619

162 &lt;212&gt; TYPE: DNA

163 &lt;213&gt; ORGANISM: Candida albicans

W--> 165 <220> FEATURE: ~~DNA~~ delete - DO NOT insert anything on the <220> line.  
 <220> is a "header" only, and stands alone.

## RAW SEQUENCE LISTING

DATE: 10/29/2001

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TIME: 15:02:39

Input Set : A:\JAB-1430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

Explanation  
goes on  
line  
2237

W--> 166 ~~221~~ NAME/KEY: n = any nucleotide >2237 <←

168 <400> SEQUENCE: 6

169 aactaatttg tttaaacatc aataccaaga agttttttaca attcaatccc acatacacca 60

170 ttaattatga attctgaaaa gattattgaa gttatcattg ctatttttctt accaccagta 120

171 gctgtgttta tgaaatgtgg tgccactacc ccattatgga ttaacttggt attatgtatc 180

172 tttattttggt tccctgctat cttacatgcc ttatacgttg tgttgaaaaga ttaaacaac 240

173 accagagatt tactgcttga tgaattgatt actccaaaga gttgtgacta gttcccagtg 300

174 tgtttttttt gccttccaac tttctttttac atttttccat tactaccact gtcttccccc 360

175 ctatttttgca gagttttcaa aatttatcca aaacatgtta gtcattaaac catattatta 420

176 taattattct tttttgtatt tttttccctt aaaacacggt aattttattaa tcgtttcgtt 480

177 gtttggattt ttattttttt gtatttatca atttgaatat atatctatac atgaatttat 540

W--> 178 tatccattgt accaattggt aaaacatttt gttagttttt tggtactagt ataaaannat 600

W--> 179 aataaaagtt tanttcaac 619

181 <210> SEQ ID NO: 7

182 <211> LENGTH: 2319

183 <212> TYPE: DNA

184 <213> ORGANISM: Candida albicans

186 <400> SEQUENCE: 7

187 atgacattag ggttcgataa attcataagc aagggtcagca ctcatagacg tcaatctgaa 60

188 ccatcaatct tggaaatcgc agccaccaat tctcaaaata aatcgagaag gctaagtatg 120

189 gataatggtc attgttatgt tctgtgaatca actaataatc atcatcattt aaataccgtc 180

190 gttgataaatt tacgacagcg tgccgggatcg ttttcattta tttcacatca ccataatcac 240

191 catcagaata gtcacgataa ttatactgtc gatccctta catcaaacgg agcacgaatt 300

192 tcccgatcac gttcacgttc caaatcagtt gggcacggag aagcaatatc accagcgtat 360

193 ttttccaaga ataaaaccaa agatttagtg aaacaggaaa cagcacatat cattctgaag 420

194 aaattactca acatgttaca agatttggtt ttacaaaacc ctattgcatt gaaaacaata 480

195 tcacaagggt cagaatcaaa gttttgtaaa atctacgtgt ctaacactaa taattgtatt 540

196 tacttaccag cagcaagttc aacaagtttc acttatgaag atgatgaaaa tggcggcgtt 600

197 ataattgtcg aagatagaaa tgatgaaatg ccaacagcag ttaataacaa tactttgtca 660

198 atggatagta taaatcattc agagactgat ttccctggatt ctccaccacc tccagattta 720

199 ttttctaaaa tgaaatcatt ccattcacca aattacttga cttcaaaaat cgattctgaa 780

200 tgtccaattc cacatacatt tgctgtgatt gttgaattaa ccaaggactc tttgattatt 840

201 aaagatcttc atttccaatt tcagtcatta actaccattt tatggccaac tggggatgca 900

202 tataatcgga ctcatgccaa ggagaaattt accattggga atatggaatg gcgtacatct 960

203 ttaagcgacg ccgactatta tatcaatagt tctaattcca acgatgttaa gctgaaaaac 1020

204 ttgggtcctg aagatcttat taatcgaact agagaatata aattaatcga tattgaagaa 1080

205 ccaaacaatt catcaaacag tttactggat gatgacatgg atattaataa tattacgtcg 1140

206 ccattatcaa cgtcaccaac atcaagttca acttcaacaa attcaacctc caactcattg 1200

207 ggttcagatt catataaagc tggctctttat gtatttttat taccaatctt attgccagaa 1260

208 catattcctg cttccattgt ttctattaat gtttcattgg ctcatacatt actggttgaa 1320

209 tgcaataaat atactgataa gttgaatcgg aaatcaaaaag tatcagcatc gtacaattta 1380

210 cctatgggtc gtaactccacc aaacattgggt aattccattg ctgataagcc aatttatgtt 1440

211 aataggattt ggaatgatgc cgtacattat attataactt tccccgcaa atatgttact 1500

212 ttgggttggt aacacatgat aaatgtgaaa ttactgccca tggtgaaaaga tgtggttatc 1560

213 aagcgtatta aatttaattgt attggagaga ataacttatg tttccaaaaa tttatcacga 1620

214 gaatatgatt atgatagtga agacccttat tgtattcatc cagtttctaa agaaaataaa 1680

215 gtacgtgaac gtgttgtgtc gttatatgaa ttgaaaacga aggcaaaaca atcttctggt 1740

216 ggacatcttg aagcttataa acaagaagtt atgaaatgtc cggaaaaataa ctttttattt 1800

217 tcttgttatg aggttgaaaa tgataataat aacggcaacg gcaacggcaa cggcaacgga 1860

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/857,372A

DATE: 10/29/2001

TIME: 15:02:39

Input Set : A:\JAB-1430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

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218 aacaagaacg ttaaacaataa gaataaagat caaccaatga ttgctacacc tttagatatc 1920
219 aatgttttctt taccattttt aactactatg tctgatagtt taattatgac atcagccata 1980
220 gaagaagaag gttcagatct gcctcatata tcaagaagag ggtcggcagt gaggatgact 2040
221 gataataata ctacccaag taacaataac cctttatctc ctttttggg agcagtggaa 2100
222 actaatggtg ctagtataaa tgaaattggt gatcatatcat tattccctga ttctaatttt 2160
223 cgacatatgt aaattaaaca tgcattacaa gttacattta ggatttctaa accggatctg 2220
224 gataataaaa tgcattacaa tgaagtgggt attgataccc ccatcgtttt acttagttca 2280
225 aaatgtcaag aagatttctc tctccttat agttctgta 2319

```

227 &lt;210&gt; SEQ ID NO: 8

228 &lt;211&gt; LENGTH: 255

229 &lt;212&gt; TYPE: DNA

230 &lt;213&gt; ORGANISM: Candida albicans

232 &lt;400&gt; SEQUENCE: 8

```

233 aacgttcgtg caaaaggcta tactggtgat atccacgcag atgaagagca agtttaatca 60
234 actctttgtc aattaatgct gtacttggtt tcattttatt tgctggcatt taaagaatac 120
235 ccatagttca gaaaataaaa ttgaaaaatt taaaaaaaaa cgcaatatca ttcatttttt 180
236 ttgttttttt gacaataata ttaatatgta gttaccaatg ttttagatt ttatatgttt 240
237 tgaaaaaata gtttg 255

```

239 &lt;210&gt; SEQ ID NO: 9

240 &lt;211&gt; LENGTH: 119

241 &lt;212&gt; TYPE: DNA

242 &lt;213&gt; ORGANISM: Candida albicans

W--> 244 <220> FEATURE: ~~DNA~~ *delete*W--> 245 <221> NAME/KEY: ~~n = any nucleotide~~ *>L2237<*

247 &lt;400&gt; SEQUENCE: 9

OK-> 248 aaccttacaa tcattatacc aactatcaaa atcataagac tcttnaactt ctgtttttga 60  
 249 tagttggtat aatgatttat gtattatctt aattcattat tattagtttc ggtcacaaa 119

251 &lt;210&gt; SEQ ID NO: 10

252 &lt;211&gt; LENGTH: 60

253 &lt;212&gt; TYPE: PRT

254 &lt;213&gt; ORGANISM: Candida albicans

256 &lt;400&gt; SEQUENCE: 10

```

257 Met Ile Thr Asp Glu Gln Leu Asn Thr Ile Ala Leu Thr Phe Gly Phe
258 1 5 10 15
260 Ala Ser Ile Ile Leu Ile Ile Ile Tyr His Ala Ile Ser Thr Asn Val
261 20 25 30
263 His Lys Leu Glu Asp Glu Thr Pro Ser Ser Ser Phe Thr Arg Thr Asn
264 35 40 45
266 Thr Thr Glu Thr Thr Val Ala Ser Lys Lys Lys Lys
267 50 55 60

```

270 &lt;210&gt; SEQ ID NO: 11

271 &lt;211&gt; LENGTH: 426

272 &lt;212&gt; TYPE: PRT

273 &lt;213&gt; ORGANISM: Candida albicans

275 &lt;400&gt; SEQUENCE: 11

```

276 Met Leu Lys Thr Thr Gln Thr Leu Arg Leu Thr Gly Lys Ala Phe
277 1 5 10 15
279 Pro Lys Val Arg Pro Ala Leu Ile Arg Thr Tyr Ala Ala Phe Asp Arg
280 20 25 30

```

## VERIFICATION SUMMARY

DATE: 10/29/2001

PATENT APPLICATION: US/09/857,372A

TIME: 15:02:40

Input Set : A:\JAB-I430.txt

Output Set: N:\CRF3\10292001\I857372A.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:165 M:256 W: Invalid Numeric Header Field, <220> has non-blank data  
L:166 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6  
L:178 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6  
L:178 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:6  
L:178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:179 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6  
L:179 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:6  
L:179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:244 M:256 W: Invalid Numeric Header Field, <220> has non-blank data  
L:245 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9  
L:248 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9  
L:248 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9  
L:248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  
L:787 M:283 W: Missing Blank Line separator, <220> field identifier